Viewing pembrolizumab_abb2_1FH-clashlist.txt

Duke Biochemistry
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```
Using nuclear cloud x-H distances and vdW radii

Using input model H/D atoms...

Bad Clashes >= 0.4 Angstrom:
 L  43  GLN  HB2  L  53  LEU HD11 :0.520
 L  24  ARG  HE   L  86  ASP  CG  :0.459
clashscore = 0.57
clashscore (B factor cutoff = 40) = 0.567215
```

About MolProbity | Website for the Richardson Lab | Using nuclear x-H | Internal reference 4.5.2